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gcc TAA

```

Fig. 1

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1 50
EEASGKVVQD IMPCLHFVKG EEKEPSKESS SGTKKLSEEV KTTEQKREAC

51 100
KCIVRATKGI SGIKNELVAE VPKKCDIKTT LPPITADFDC SKIQSTIFRG

101 150
YYGFQETSGT MVRALMPCLP FVQGKEKEPS KGSSSGAKRL DGETKTGPQR

151 200
VHACECIQTA MKTYSIDIDGK LVSEVPKHCG IVDSKLPPID VNMDCKTVGV

201 243
VPRQPQLPVS LRHGPVTGPS DPAHKARLER PQIRVPPPAP EKA

Fig.2

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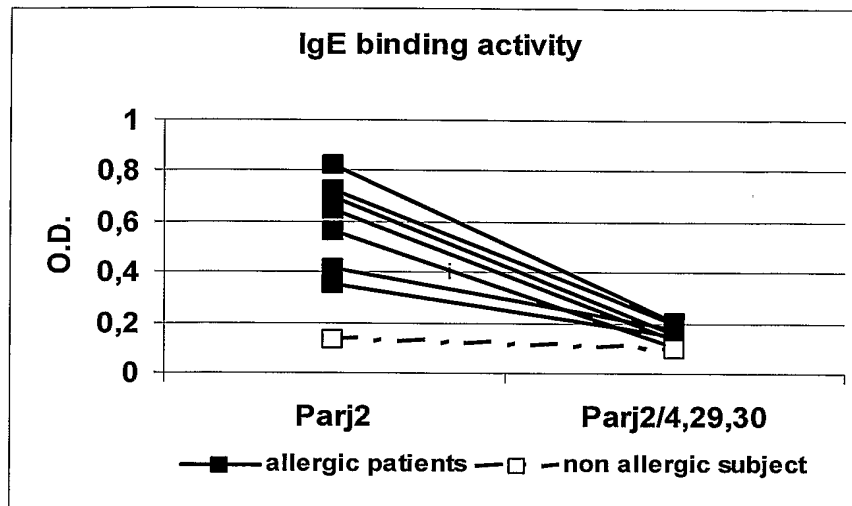


FIG. 3

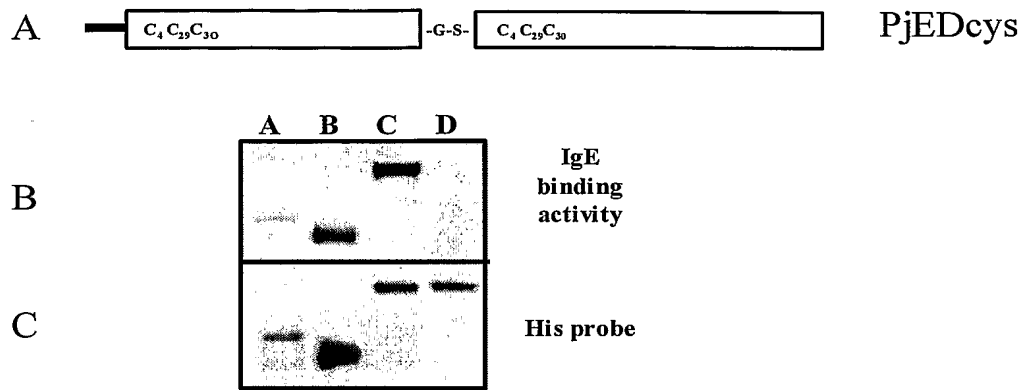


Fig 4

Binding inhibition
ELISA

	Dimer W.T	PjEDcys
Patient 1	68%	7%
Patient 2	56%	3,5%
Patient 3	62%	10%
Patient 4	62%	8%
Patient 5	62%	8%

Fig. 5

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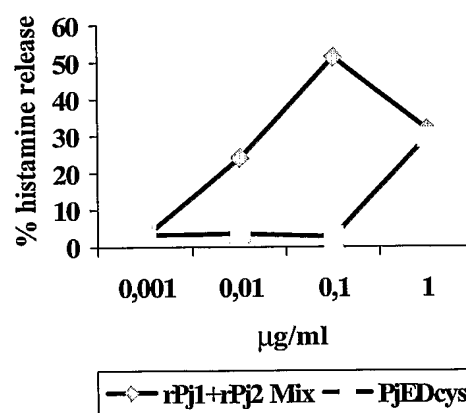
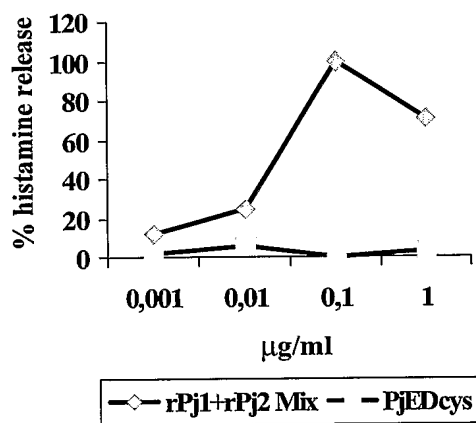
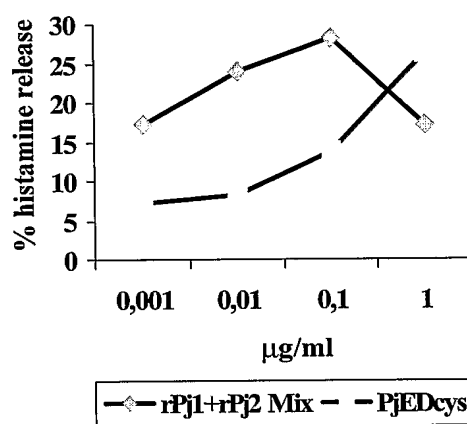
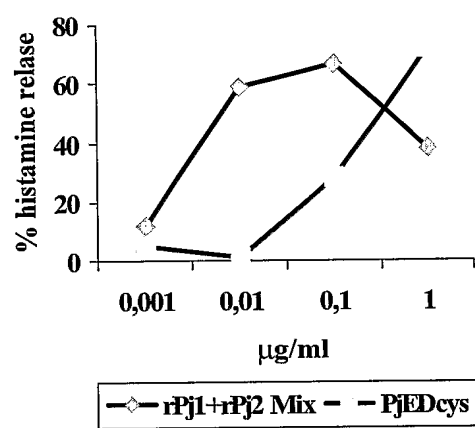


Fig. 6

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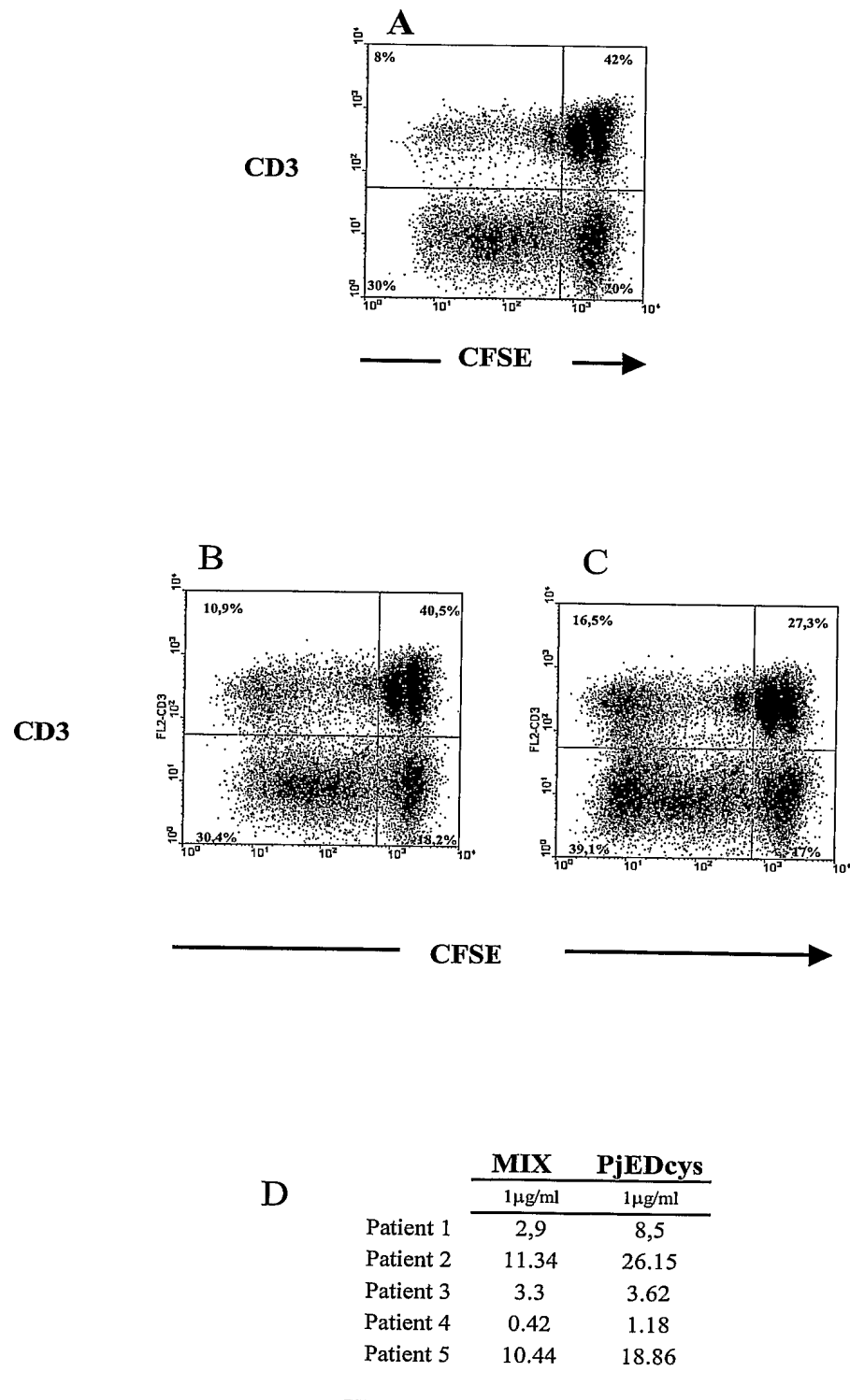


Fig 7

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**% of human IgE
binding inhibition**

<i>P. judaica</i> extract	Expj	rParj1	rParj2	rParj1 rParj2
Patient 1	94	10	80	80
Patient 2	93	53	74	74
Patient 3	90	35	37	80
Patient 4	98	59	85	86
Patient 5	100	57	29	67
Patient 6	91	29	65	69
Patient 7	94	38	58	79
Average	94,3	40,1	61,1	76,5

Fig. 8

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**% of human IgE
binding inhibition**

tested Protein	% of inhibition
rParj 1	95%
PjA	16%
PjB	85%
PjC	14%
PjD	15%

Fig. 9